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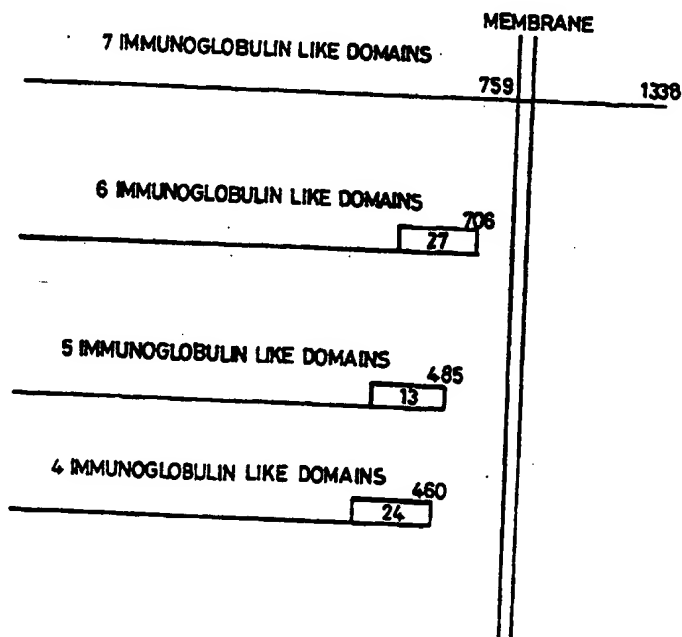
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(54) Title: FLT-4 (FMS-LIKE TYROSINE-KINASE), FLT-15, VARIANTS THEREOF USES AS GROWTH FACTOR INHIBITORS



## (57) Abstract

Disclosed is an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains, together with pharmaceutical compositions comprising the polypeptide, and various uses thereof.

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FLT-4 (fms-like Tyrosine kinase), FLT-15, variants thereof used as growth factor inhibitors

### Field of the Invention

This invention relates to substances which inhibit growth factors, in particular, vascular endothelial growth factor (VEGF), methods of inhibiting growth factors and of treating tumours and regulating fertility.

### Background of the Invention

A considerable number of human growth factors are now known, many of which have been at least partly characterised. Among them is vascular endothelial growth factor (VEGF), which has been identified in several tissues (Gospodarowicz *et al.*, 1989 PNAS 86, 7311-7315; Conn *et al.*, 1990 PNAS 87, 2628-2632; Tischer *et al.*, 1991 J. Biol. Chem. 266, 11947-11954). As its name suggests, this growth factor is a highly specific mitogen for endothelial cells and is greatly involved in angiogenesis. VEGF is a homodimeric glycoprotein of two 23kDa subunits exhibiting sequence homology with platelet-derived growth factor A and B chains and placenta growth factor.

The homologous tyrosine kinase receptors fms-like tyrosine kinase receptor (FLT) and kinase insert domain-containing receptor (KDR) function as high-affinity VEGF receptors (de Vries *et al.*, 1992 Science 255, 989-991; Terman *et al.*, 1992 Biochem. Biophys. Res. Commun. 187, 1579-1586). Both FLT and KDR are membrane-spanning receptors that each contain seven immunoglobulin-like domains in the extracellular ligand-binding region, an intracellular tyrosine kinase domain and a transmembrane domain. The transmembrane domain serves to anchor the receptor in the cell membrane of the cells in which it is expressed.

A number of membrane-bound receptor molecules have been found to exist in truncated soluble forms, generated either by proteolytic processing or by alternative splicing of

mRNA. Recently, Kendall & Thomas (1993 PNAS 90, 10,705-10,709, and WO94/21679) described the discovery of a soluble form of FLT receptor (sFLT) generated by alternative splicing.

Essentially, Kendall & Thomas screened a human umbilical vein endothelial cell (HUVEC) cDNA library with one probe specific for the 3' end of the flt coding region (encoding the intracellular tyrosine kinase domain) and with another probe specific for the 5' flt coding portion (encoding one of the extracellular N terminal domains). Clones which hybridised with the 5' specific probe but not with the 3' specific probe were selected for further study. In this way, a clone was isolated which encoded a soluble FLT polypeptide lacking the transmembrane domain and the intracellular domain. The truncation resulted from "readthrough" to an intronic termination codon. It was suggested by Kendall & Thomas that the soluble receptor could act as an efficient specific antagonist of VEGF *in vivo*.

The present invention is based on the discovery of further soluble variants of FLT, the existence of which was not predicted by the teaching of Kendall & Thomas.

#### Summary of the Invention

In a first aspect the invention provides an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains. Preferably, the altered FLT polypeptide comprises four or fewer complete Ig-like domains. The altered soluble FLT polypeptide inhibits VEGF by preventing it binding to its natural receptors, flt and KDR, present on the surface of target cells. Surprisingly, such truncated forms, lacking a major extracellular portion of the molecule, are believed to retain affinity for VEGF.

The term "soluble" as used herein is intended to refer to altered forms of the FLT polypeptide which do not comprise a transmembrane domain and thus generally do not become associated with the cell membrane of cells in which the molecule is expressed. In particular, the invention provides soluble altered forms of the FLT polypeptide

consisting substantially of four or five complete immunoglobulin-like domains.

In a particular embodiment the invention provides an altered, soluble form of FLT having at its C-terminus a region substantially having the amino acid sequence of the sequences termed FLT4 or FLT15 shown in Figure 5, or a functional equivalent thereof. The term "functional equivalent" as used above is intended to include those polypeptides which have substantially the same deletions as the polypeptides encoded by FLT4 or FLT15 (with respect to the unaltered full length FLT molecule), but which may also have other deletions, additions or substitutions, (in particular conservative substitutions), and which retain an inhibitory effect for VEGF.

Preferably the polypeptide will also comprise, at its N-terminus, the amino acid sequence substantially corresponding to the equivalent portion of the unaltered wild-type FLT polypeptide. Conveniently, polypeptides in accordance with the invention will comprise around 400 to 500 amino acid residues, preferably around 480 amino acid residues, most preferably between 480 and 440 amino acid residues of the wild type FLT sequence.

Preferably the polypeptides of the invention arise by alternative splicing of mRNA or by proteolytic processing of a mature polypeptide, although it will be apparent to those skilled in the art that the polypeptide could be encoded by a nucleic acid derived, at least in part, by recombinant DNA technology.

In a further aspect the invention provides a nucleic acid sequence encoding a polypeptide in accordance with the invention. In a particular embodiment the invention provides a nucleic acid comprising the sequence of nucleotides inserted at position 1655 of the FLT 4 sequence shown in Figure 3 or the sequence of nucleotides inserted at position 1555 of the FLT 15 sequence shown in Figure 3, or a functional equivalent thereof. Examples of functionally equivalent nucleic acids include those sequences which encode substantially the same polypeptide as those encoded by FLT4 or FLT15 but which differ in nucleotide sequence as a result of the degeneracy of the genetic code. It will be apparent to those skilled in the art that the portion of the inserted nucleotide sequence in FLT4 and FLT15 occurring after the premature termination codon could be omitted without affecting the characteristics of the encoded polypeptide. Accordingly, nucleic acid molecules without

such sequences are also regarded as functionally equivalent for the purposes of the present invention.

Conveniently, the nucleic acid will substantially comprise the nucleotide sequence of FLT4 or FLT15 shown in Figure 3, together with the nucleotide sequence encoding the N-terminus of unaltered, wild-type FLT. Advantageously, the nucleic acid will be obtainable by means of PCR amplification from a sample of human cells. Desirably, the nucleic acid will be obtainable by means of PCR using primers intended to hybridise to non-conserved regions of the FLT molecule. Conveniently, the nucleic acid sequence will be obtainable by use of PCR primers designed to hybridise to the regions of the FLT sequence shown underlined in Figure 3, or immediately adjacent thereto. In particular, the PCR primers will conveniently have substantially the sequence: 5'-GCAAGGTGTGACTTTGTTC-3' and 5'-AGGATTCTTCCCCTGTGTA-3'.

In another aspect, the invention provides a method of inhibiting VEGF *in vitro*, comprising adding an effective amount of the polypeptide defined above. It may also be desirable to inhibit VEGF in a human subject. Thus the invention provides a method of inhibiting VEGF in a human subject, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. In particular, VEGF provides a mitogenic stimulus (particularly involved in angiogenesis), so inhibition of VEGF would be expected to provide therapeutic effects in the treatment of tumours or disorders involving inappropriate neovascularisation.

In particular the invention provides for a method of treating tumours or diseases involving inappropriate neovascularisation, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. Suitable diseases which might be amenable to treatment include ovarian cancer and ovarian hyperstimulation (Boocock *et al.*, 1995 J. Natl. Cancer Inst. 87, 506-516).

Furthermore, it has been conclusively demonstrated that FLT is expressed by trophoblasts and cells from ovarian and endometrial tissues (Charnock-Jones *et al.*, 1994 Biology of Reproduction 51, 524-530), which clearly suggests a role for VEGF in the growth and

differentiation of trophoblasts during implantation.

Thus, in particular, the invention provides a method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance.

It will be appreciated by those skilled in the art that the identification of FLT on the surface of trophoblasts and endometrial cells also provides a number of possible methods of regulating fertility. For example, the growth of trophoblasts is essential for successful implantation of the embryo. Inhibition of trophoblast growth thus provides a method of contraception or contragestion.

Thus in a further aspect the invention provides a method of regulating the fertility of a human female, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. An "effective amount" of the polypeptide is an amount sufficient to substantially block the stimulus of VEGF on trophoblasts and/or endometrial cells. Typically, the method will result in reducing the fertility of the female.

Moreover, it might be possible to identify agents which can enhance the effect of VEGF on trophoblasts, and thereby improve the probability of successful implantation, either in assisted or spontaneous cycles. Candidates for such VEGF-enhancing agents would include anti-sense equivalents of the nucleic acid sequences encoding the truncated FLT polypeptides of the invention. It will be apparent to those skilled in the art that these could be used to improve the fertility of a human female.

In a further aspect the invention provides a pharmaceutical composition comprising the polypeptide defined above, together with a physiologically acceptable carrier substance. The composition could be used in vivo any one of the methods defined above. In yet another aspect the invention provides for the use of a polypeptide in accordance with the invention in the preparation of a therapeutic composition for the treatment of tumours and

diseases involving inappropriate neovascularisation. Examples of such conditions and diseases are detailed, inter alia, in WO94/10202 and WO94/21679. The invention also includes within its scope a method of making a pharmaceutical composition, comprising mixing the polypeptide defined above together with a physiologically acceptable carrier substance.

The invention will now be described by way of the following illustrative examples and with reference to the drawings, of which:

Figure 1 shows an amino acid multiple alignment of closely related tyrosine kinase receptors (flt, fms and kit, "kit" being another name for KDR);

Figure 2 shows typical results of agarose gel electrophoresis demonstrating the existence of alternatively-spliced flt-coding sequences in various tissue samples;

Figure 3 shows the nucleotide sequence of the 3' region of the sequences encoding full length VEGF receptors (FLT and the related receptor KDR), together with two sequences, FLT4 and FLT15, which encode polypeptides according to the invention;

Figure 4 is a schematic representation of wild type and mutant FLT molecules; and

Figure 5 shows the C terminal amino acid sequences of two polypeptides in accordance with the invention.

#### Example

Expression of FLT, the VEGF receptor, was investigated in cell lines derived from human trophoblast-like and ovarian and endometrial carcinomas. The trophoblast-like (choriocarcinoma) cell line used was BeWo (obtained from the American Type Culture Collection, Rockville MD, USA). The endometrial carcinoma cell lines were Ishikawa (obtained from Professor M Nishide, University of Tsukuba, Japan), and HEC 1-A and HEC 1-B (from ATCC, USA). The ovarian cancer cell lines were 7, 17R, 25, 25R and 35. These were all shown to be of epithelial origin and had been established in culture



for 10-30 passages. Cell lines 17R and 25R were derived after chemotherapy and subsequent relapse (line 25R originating from the same patient as line 25).

BeWo cells were grown in Ham's F12, according to ATCC recommendations. Endometrial carcinoma lines were grown in McCoy's medium (ICN Flow Laboratories, Irvine, UK) with 10% foetal calf serum (ICN Flow) plus 2mM L-glutamine (ICN Flow) and 50U/ml and 50mg/ml penicillin/streptomycin (ICN Flow).

It was decided to investigate expression of FLT in these cell lines and normal tissues by performing PCT and *in situ* hybridization. It was therefore necessary to construct suitable oligonucleotide primers and probes.

To help design appropriate primers, a protein multiple alignment of closely related tyrosine kinase receptors (FLT, FMS and KIT) was constructed (shown in Figure 1) using the computer program "pileup". This revealed regions of divergent sequence among this family of receptors. The regions chosen for primer design are shown with double underlining in Figure 1. The following nested PCR primers were then synthesized based on these protein sequences:

- A) 5' GCAAGGTGTGACTTTTGTTC 3'
- B) 5' GCGCTCGAGAGCATCACTCAG 3'
- C) 5' GCGCGGCCGCAGTAAAATCCA 3'
- D) 5' AGGATTTCCTCCCCTGTGTA 3'

The underlined portions of these oligonucleotides are the regions which hybridise to the flt cDNA sequence. The other nucleotides were added to facilitate directional cloning. The cycles used were: first round with primers A and D [95°C 30 seconds, 55°C 30 seconds, 72°C 30 seconds] x 25; second round with primers B and C: [95°C 30 seconds, 44°C 30 seconds, 72°C 30 seconds] x 2 [95°C 30 seconds, 65°C 30 seconds, 72°C 30 seconds] x 25. The internal primers B and C had sites for the restriction enzymes Xho I and Eag I respectively at their 5' ends to permit directional cloning of the products.

It was found that certain tissues gave rise to PCR amplification products of notably larger size (as judged by agarose gel electrophoresis) than observed for the full length FLT cDNA product. Typical results are shown in Figure 2.

PCR products obtained using the nested set of primers A-D were run out on a gel. Lanes 1-3 are products obtained from primary tissue samples of the ovarian carcinomas designated 17, 17R and 25R. Lanes 4 to 7 are products obtained from cell lines established from the ovarian carcinomas 7, 17R, 25 and 25R. Lanes 8 to 10 are the cell lines HEC 1-A, HEC 1-B and Ishikawa respectively. Lane 11 contains products from HUVECs.

The standard size band was of the expected size (around 285bp) and was found to be identical to the 3' end of the published flt sequence (Shibuya *et al.*, 1990 Oncogene 5, 519-524). However it can be clearly seen that in addition to the full length flt cDNA PCR-amplified product, in lanes 2 (17R, primary tissue) and 4 (7, cell line) are larger bands of approximately 360bp. A faint band of similar size was also apparent in lane 5 (17R, cell line) but is not clearly seen when the gel photograph is reproduced. These larger bands were extracted from the gel by known techniques and subcloned into the plasmid vector pBluescript II KS and then subjected to sequence analysis using the dideoxynucleotide sequencing method (Sanger *et al.*, 1977 PNAS 71, 5463-5467).

Sequencing of five independent clones (Boocock *et al.*, 1995 J. Natl. Cancer Inst. 87, 506-516) revealed that each contained one of two novel insertions within the published flt sequence, in the region between the primers. Three of these clones (termed FLT5, FLT15 and FLT16) contained an 85bp insertion at about position 1555, whilst two other clones (FLT13 & FLT14) contained a 65bp insertion at about position 1665 (see Figure 3, numbering based on that of Shibuya *et al.*, 1990 cited above). The insertions account for the larger band size of the PCR products. However, both insertions contain an in-frame termination codon, so that corresponding full length RNAs would encode soluble, truncated receptor variants comprising the first five immunoglobulin-like domains of the extracellular region, up to amino acid 517 or 553, with either 24 or 14 (of which 13 are additional) unrelated amino acids at the C-terminus.

Although these variant flt clones were derived from partial cDNAs encoding only amino acids 503 onward, PCR products of the sizes predicted for corresponding full length cDNAs were amplified from cDNA derived from HUVEC cells, human chorion and ovarian carcinoma cell line 7, using primers specific for each of the novel insertions together with a primer binding just 5' of the initiating ATG (data not shown).

Figure 4 is a schematic representation of various FLT receptor molecules. At the top, (a) shows the wild type, full length FLT receptor molecule, (b) represents the truncated version described by Kendall & Thomas, (c) represents the polypeptide encoded by FLT4 and (d) represents the polypeptide encoded by FLT15. The numerals at the right show the number of amino acids in the molecule and numerals in the boxes represent the number of amino acids present in the sFLT variants but not in the wild type molecule.

Figure 5 shows the predicted C terminal amino acid sequence of the polypeptides which would be encoded by "full length" FLT4 and FLT15 clones (i.e. clones which contained all the nucleotide sequence 5' of the primer site used to generate the actual clones). The last 14 amino acids of the FLT4 clone, and the last 24 amino acids of the FLT15 clone, are divergent from the wild type FLT sequence.

### Claims

1. An altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains.
2. A polypeptide according to claim 1, comprising four or fewer complete immunoglobulin-like domains.
3. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT4 as shown in Figure 5, or a functional equivalent thereof.
4. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT15 as shown in Figure 5, or a functional equivalent thereof.
5. A polypeptide according to any one of the preceding claims, comprising around 400 to 500 amino acid residues of the wild type FLT polypeptide.
6. A nucleic acid sequence encoding a polypeptide in accordance with any one of the preceding claims.
7. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1655 of the FLT4 sequence shown in Figure 3, or a functional equivalent thereof.
8. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1555 of the FLT15 sequence shown in Figure 3, or a functional equivalent thereof.
9. A method of inhibiting VEGF *in vitro*, comprising adding an effective amount of a polypeptide in accordance with any one of claims 1 to 5.

10. A method of inhibiting VEGF in a human subject, comprising administering an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
11. A method according to claim 10, comprising the use of a polypeptide in accordance with any one of claims 1 to 5 in the treatment of tumours or diseases involving inappropriate neovascularisation.
12. A method according to claim 11, for the treatment of ovarian cancer, ovarian hyperstimulation, or endometriosis.
13. A method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
14. A method of regulating the fertility of a human female by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
15. A pharmaceutical composition for use in the method of any one of claims 11 to 14, comprising a polypeptide in accordance with any one of claims 1 to 5, and a physiologically acceptable carrier substance.
16. A method of making a composition according to claim 15, comprising mixing a physiologically acceptable carrier substance together with a polypeptide according to any one of claims 1 to 5.

50  
 1  
 kit MRGARGAWDF LCVLLLLLLRV OTGSSQPSVS PGEPSPPSIH PGKSDLIVRV  
 fms .....M GPGVLLLLLV ATAWHGQIP VIEPSVP... ..ELVVKP  
 flt MVS YWDTGVL LCALLSCLLL TGSSSGSKLK DPESLKG... ..TOHMQA

100  
 51  
 kit GDEIRLLCTD PGEVKW.... ..TFEILD ETNENKQ... ..NEWITE  
 fms GATVTLRCVG NGSEWDGP. ....ASPHWT LYSDGSS... ..SILSTN  
 flt GQTLHLQCRG EAAHKWSLPE MVSKESERLS ITKSACGRNG KQFCSTLTIN

150  
 101  
 kit KAEATNTGKY TC....T... NKHGLSNSIY VFVRDPAKLF .....LVDRS  
 fms NATFQNTGTY RC....TEPG DPLGGSAAIH LYVKDPAFPW ....NVLAE  
 flt TAQANHTGFY SCKYLAVPTS KKKETESAII IFISDTGRPF VEMYSEIPEI

200  
 151  
 kit LYGEDNDTL VRCPLTDPEV .TNYSLKGCQ GKPLPKD.LR FIPDPKAGIM  
 fms VVFEEDQDAL LPCLLTDPVL EAGVSLVRVR GPPLMRH.TN YSFSPWHGFT  
 flt IHMIEGRELV IPCRVTSPII ..TVTLKKFP IDTLIPDGKR IIWDSRKGEI

250  
 201  
 kit IKS VKRAYHR LCLHCSVDQE GKS VLSEKFI LKVRPAFKAV PVVSVSKASY  
 fms IHRAK.FIOS ODYQCSALMG GRKVMISISR LKVQKVTPGP PALTLVPAEL  
 flt ISNAT.YKEI GLLTCEATVN GHLYKINYLT HROTNTIIDV QISTPRPVKL

300  
 251  
 kit LLREGEEFTV TCTI.KDVSS SVYSTWKREN SQT KLOEK... ..YNSWHHGD  
 fms VRIRGEAAQI VCSA.SSVDV NEDVFLQHN ..TKLAIP... ..QOSDFHNN  
 flt L..RGHTLV L NCTATTPLNT RVQMTWSYPD EKNKRASVRR RIDQNSHAN

350  
 301  
 kit FNYERQATLT ISSARVND SG VFMCYANNIF GSANVTITTE VVDKGFINI.  
 fms .RYQKVLTLN LDCVDFQHAG NYSCVASNVQ GKHSTSMFTR VVESAYLNL.  
 flt IFYS...VLT IDKMNKDKG LYTCRVRS GP SFKSVNTSVH IYDKAFITVK

400  
 351  
 kit FPMINTTVFV NDCENVDLIV EYEAFFKPEH QOWIYMNRTF TDKWEDYPKS  
 fms SSEQNLIQEV TVGEGLNLKV MVEAYPGLOG .....FNWTY LGPFSDHOPE  
 flt HRKQOVLETV AGKRSYRLSM KVKAFFSPEV V..... ..WLKD

450  
 401  
 kit ENESN..... .IRYVSELHL TRKGTGGT YTF LVS..NS DVNAALAFNV  
 fms PKLANATT KD TYRHFTLSL PRLKPSEAGR YSEFLAR..NP GGNRALTFEL  
 flt GLPATEKSAR YLTRGYSLII KDVTEEDAGN YTILLSIKQS NVEKNLTATL

Fig. 1 Sheet 1

SUBSTITUTE SHEET (RULE 26)

2/8

451

kit YVNTKPEI.. LTYDRL.... ..VN..GML QCVAAGFPEP TIDWYFCPGT 500  
 fms TLRYPPEV.. SVTWIF.... ..INGSGTL LCAASGYPOP NVTWLQCSGH  
 flt IVNVKPOIYE KAVSSFPDPA LYPLGSRQIL TCTAYGIPOP TIKWEWHPCN

501

kit EORC..... 550  
 fms TDRCD.....  
 flt HNHSEARCF CSNNEESFIL DADSNMGNRI ESITORMAI EGKNKMASTL

551

kit ..... 600  
 fms ..... SASV  
 flt VVADSRISGI YICIASNKVG TVGRNISFYI TDVPNGEHVN LEKMPTEGED

601

kit LPV..DVQTL NSSGPPF... 650  
 fms LQWDDPYPE VLSQEPF... ..GKLVVQSS  
 flt LKLSCVKNKF LYRDVWILL RTVNNRIMHY SISKQKMAIT KEHSITLNL

651

kit IDSSAFKING TVECKAYNDV G..... 700  
 fms LTVETLEHNO TYECRAHNSV G.....  
 flt IMNVSLQDSG TYACRARNVY TGEEIL OKKE ITIRDQEPY LLRNLSDHTV

701

kit ..KTSAYFNF A..... 750  
 fms ..SGSWAF.I P..... ..FKGNKEQ IHPHTLETP.  
 flt AISSSTTLDC HANGVPEPQI TWFKNNHKIQ QEPGIILGPG SSTLFIERVT

751

kit ..... 800  
 fms ..... LLI GEVIVAGMMC  
 flt EEDEGVYHCK ATNOKGSVES SAYLTVQGT S DKSNNLELITL TCTCVAATLF

801

kit IIVMILTYKY LQKPMYEVOW KVVEEINGNN YVYID..PTQ LPYDH.KWEE 850  
 fms LLLLLLLLYKY KQKPKYQVRW KIIESYEGNS YTFID..PTQ LPYNE.KWEE  
 flt WLLLTLLIRK MKRSSEIKT DYLSIIMDPD EVELDEQCER LPYDASKWEE

851

kit PRNRLSFGKT LGAGAFGKVV EATAYGLIKS DAAMTVAVKM LKPSAHLTER 900  
 fms PRNNLOFGKT LGAGAFGKVV EATAFGLGKE DAVLKVAVKM LKSTAHADK  
 flt ARERLKLKGS LGRGAFGKVV QASAFGIKKS PTCRTVAVKM LKEGATASEY

Fig. 1 Sheet 2

SUBSTITUTE SHEET (RULE 26)

3/8

901 950  
 kit EALMSELKVL SYLGNHMNIV NLLGACT.IG GPTLVITEYC CYGDLINFLR  
 fms EALMSELKIM SHLGOHENIV NLLGACT.HG GPVLVITEYC CYGDLINFLR  
 flt KALMTELKIL THIGHHLNV NLLGACTKQG GPLMVTVEYC KYGNLSNYLK

951 1000  
 kit RKROSEI... ..C...SKOE DHAEEALYKN L.....LHS KESSCSDSTN  
 fms RKAAML... ..GPSLSPGO DPEGGVYKN IHLEKKYVRR DSGFSSQGV  
 flt SKRDLEFLNK DAALHMEPKK EKMEPGLEQG KPRLDVTS SESFASSGFO

1001 1050  
 kit EYMDMKPGVS YVPTKADKR RSVRIGSYIE ROVTPAIME DELALDLEDL  
 fms TYVEMRP... ..VSTSSN... ..DSFSE QDLD....KE DGRPLELRL  
 flt EDKSL..... ..SDVEE EEDSDGEYKE ...PITMEDL

1051 1100  
 kit LSFSYQVAKG MAFLASKNCI HRDLAARNIL LTHGRITKIC DFGLARDIKN  
 fms LHFSSQVAG MAFLASKNCI HRDVAARNVL LTNGHVAKIG DFGLARDIMN  
 flt ISYSFQVARG MEFLSSRKCI HRDLAARNIL LSENNVVKIC DFGLARDIYK

1101 1150  
 kit DSNYVVKGNA RLPVKWMAPE SIFNCVYTFE SDVWSYGIFL WELFSLGSSP  
 fms DSNYIVKGNA RLPVKWMAPE SIFDCVYTVQ SDVWSYGILL WEIFSLGLNP  
 flt NPDYVRKGD TRLPLKMAPE SIFDKIYSTK SDVWSYVLL WEIFSLGSSP

1151 1200  
 kit YPGMPVDSKE YKMIKEGFRM LSPEHAPAEM YDIMKTOWDA DPLKRPTFKQ  
 fms YPGILVNSKE YKLVDGYOM AQPAFAPKNI YSIMOACWAL EPTHRPTFQQ  
 flt YPGVQMEDF CSRLREGMRM RAPEYSTPEI YQIMLDOWR DPKERPRFAE

1201 1250  
 kit IV....QLIE KOISES.TNH I.....Y SNLANCSPNR QKPVDHHSVR  
 fms IC....SFLQ EQAQEDRRER D.....Y TNLPSRSS. ....GGSGS  
 flt LVEKLGDLLO ANVOQDGKDY IPINAILTGN SGTYSTPAF SEDFFKESIS

1251 1300  
 kit INSVGSTASS SOP.....L LVHDDV.....  
 fms SSSELEEESS SEH.....L TCCEQGDIAQ PLLQPNNYOF C.....  
 flt APKFNSGSSD DRYVNAFKF MSLERIKTFE ELLPNATSMF DDYQGDSTL

1301 1350  
 kit .....  
 fms .....  
 flt LASPMLKRFT WTDSKPKASL KIDLRVTSKS KESGLSDVSR PSFCHSSCGH

1351 1389  
 kit .....  
 fms .....  
 flt VSEGKRRETY DHAELERKIA CCSPPPDYNS VVLYSTRPI

Fig.1 Sheet 3

SUBSTITUTE SHEET (RULE 26)



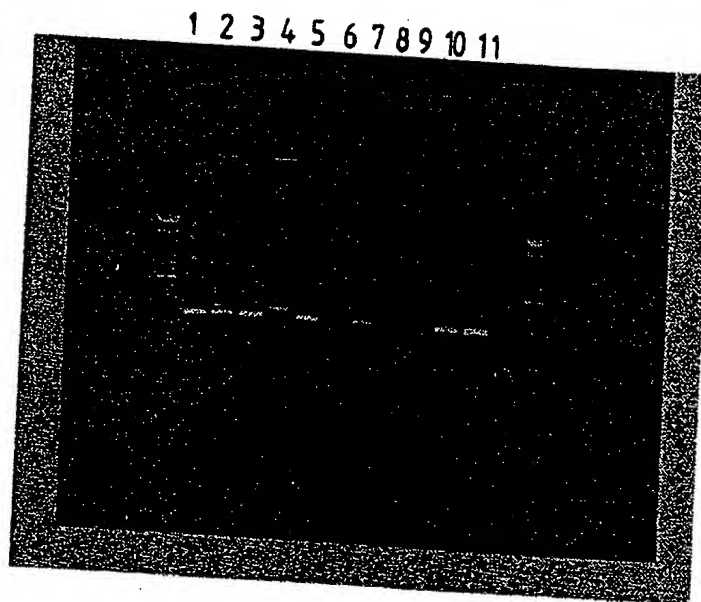


Fig. 2

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1410  
 KDR AGAGTGCGCC AACGAGCCCA GCCAAGCTGT CTCAGTGACA AACCCATACC  
 FLT ACCCCTGTAA CCATAACATT CCGAAGCAAG GTGTGACTTT TGTTCGAATA

1460  
 KDR CTTGTGAAGA ATGGAGAAGT GTGGAGGACT TCCAGGGAGG AAATAAAATT  
 FLT ATGAAGAGTC CTTTATCCTG GATGCTGACA GCAACATGGG AACAGAAATT

1510  
 KDR GAAGTTAATA AAAATCAATT TGCTCTAATT GAAGGAAAAA ACAA-----  
 FLT GAGAGCATCA CTCAGCGCAT GGCAATAATA GAAGGAAAGA ATAAG-----  
 FLT4 GAGAGCATCA CTCAGCGCAT GGCAATAATA GAAGGAAAGA ATAAG-----  
 FLT15 GAGAGCATCA CTCAGCGCAT GGCAATAATA GAAGGAAAGA ATAAGCTTCC

KDR -----  
 FLT -----  
 FLT4 -----  
 FLT15 ACCAGCTGAC AGTTCTTTCA TGTGTCACC TACAAGCTTC TCTTCCAAC

1555  
 KDR ----- CTGTAAGTAC CCTTGTTATC  
 FLT ----- ATGGCTAGCA CCTTGGTTGT  
 FLT4 ----- ATGGCTAGCA CCTTGGTTGT  
 FLT15 ACTTCCATTT CCTTCCGTGA CTCTAAACGG ATGGCTAGCA CCTTGGTTGT

1575  
 KDR CAAGCGGCAA ATGTGTCAGC TTTGTACAAA TGTGAAGCGG TCAACAAAGT  
 FLT GGCTGACTCT AGAATTTCTG GAATCTACAT TTGCATAGCT TCCAATAAAG  
 FLT4 GGCTGACTCT AGAATTTCTG GAATCTACAT TTGCATAGCT TCCAATAAAG  
 FLT15 GGCTGACTCT AGAATTTCTG GAATCTACAT TTGCATAGCT TCCAATAAAG

1625  
 KDR CGGGAGAGGA GAGAGGGTGA TCTCCTTCCA CGTGACCAGG -----  
 FLT TTGGGACTGT GGAAGAAAC ATAAGCTTTT ATATCACAGA -----  
 FLT4 TTGGGACTGT GGAAGAAAC ATAAGCTTTT ATATCACAGA ATTGTCAAAC  
 FLT15 TTGGGACTGT GGAAGAAAC ATAAGCTTTT ATATCACAGA -----

KDR -----  
 FLT -----  
 FLT4 TTTGAGTGCC TTCATCCTTG CTCTCAGGAA TAGAACTCTA CCTCATCGGA  
 FLT15 -----

Fig. 3 Sheet 1

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1665  
 KDR -----GGTCC T---GAAATT ACTTTGCAAC CTGACATGCA GCCCACTGAG  
 FLT -----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA  
 FLT4 TCTCATGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA  
 FLT15 -----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA

1710  
 KDR CAGGAGAGCG TGTCTTTGTG GTGCACTGCA GACAGATCTA CGTTTGAGAA  
 FLT AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG

FLT4 AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG  
 FLT15 AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG

1760  
 KDR CCTCACATGG TACAAGCTTG GCCCACAGCC TCTGCCAATC CATGTGGGAG  
 FLT ACGTTACTTG GATTTTACTG CGGACAGTTA ATAACAGAAC AATGCACTAC  
 FLT4 ACGTTACTTG GATTTTACTG CGG  
 FLT15 ACGTTACTTG GATTTTACTG CGG

1810  
 FLT AGTATTAGCA AGCAAAAAAT GGCCATCACT AAGGAGCACT CCATCACTCT

1860  
 FLT TAATCTTACC ATCATGAATG TTTCCCTGCA AGATTGAGGC ACCTATGCCT

1910  
 FLT GCAGAGCCAG GAATGTATAC ACAGGGGAAG AAATCCTCCA GAAGAAAGAA

Fig. 3 Sheet 2

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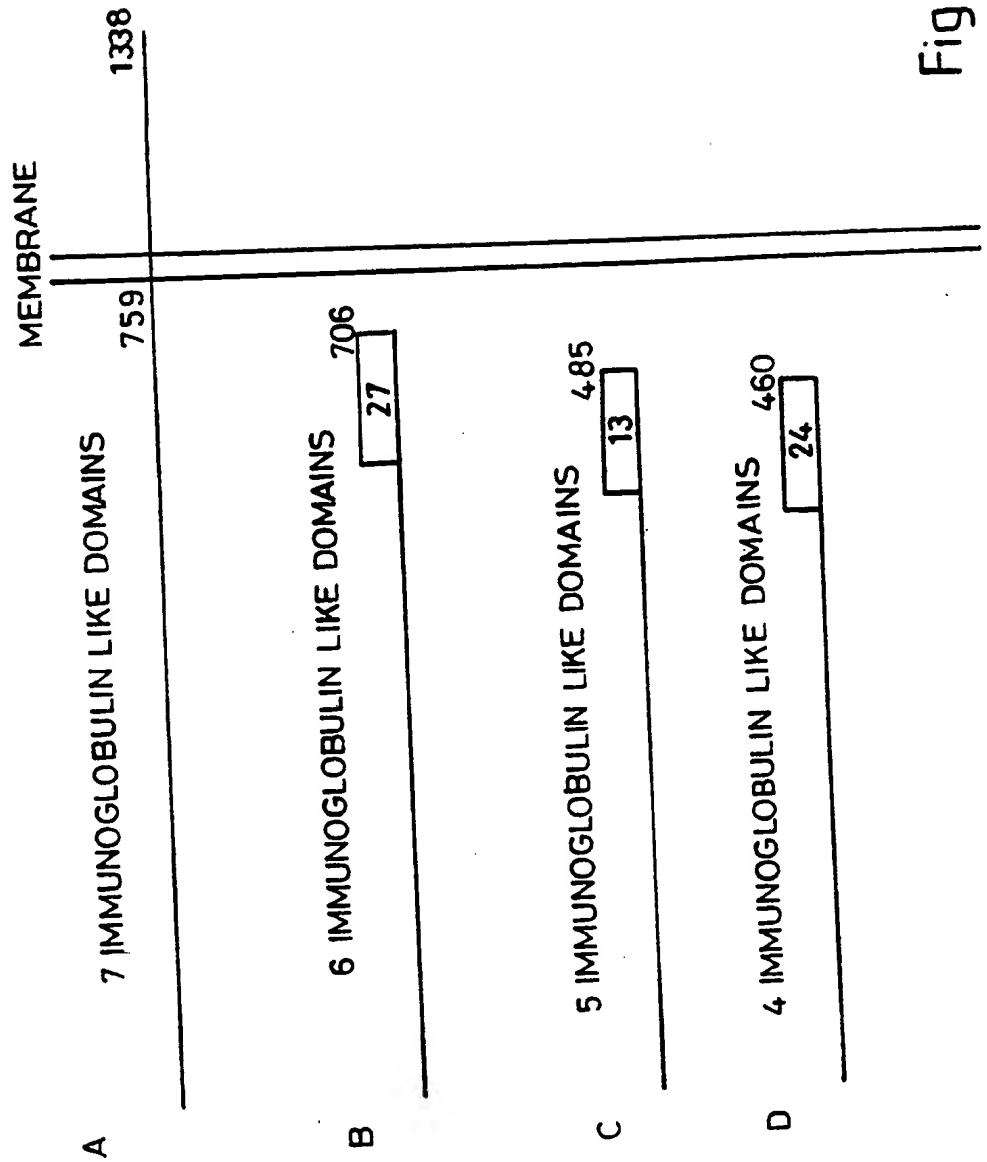


Fig. 4

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FLT4

1 ESITQRMAL EGKNKMASTL VVADSRISGI YICIASNKVG TVGRNISFYI  
51 TELSNFECLH PCSQE\*

FLT15

1 ESITQRMAL EGKNKLPPAD SSFMLPPTSF SSNYFHFLP\*

Fig. 5

## INTERNATIONAL SEARCH REPORT

International Application No

PCT/GB 95/01213

A. CLASSIFICATION OF SUBJECT MATTER  
 IPC 6 C12N15/12 C07K14/71 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	CRIT REV ONCOG, 1993, 4 (6) P595-613, UNITED STATES, ROSNET O ET AL 'Hematopoietic receptors of class III receptor-type tyrosine kinases.' see the whole document ---	1,2,9-16
Y	ONCOGENE, vol. 8, no. 11, November 1993 ENGLAND, pages 2931-2937, PAJUSOLA, K. ET AL.; 'Two human FLT4 receptor tyrosine kinase isoforms with distinct carboxy terminal tails are produced by alternative processing of primary transcripts' see the whole document --- -/--	1,2,9-16

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

## \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

\*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

\*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

\*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

\*G\* document member of the same patent family

Date of the actual completion of the international search

17 October 1995

Date of mailing of the international search report

08. 11. 95

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Authorized officer

Nauche, S

# INTERNATIONAL SEARCH REPORT

International Application No  
PCT/GB 95/01213

## C(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	ONCOGENE, AUG 1993, 8 (8) P2293-8, ENGLAND, FINNERTY H ET AL 'Molecular cloning of murine FLT and FLT4.' see the whole document ---	1,2,9-16
Y	WO,A,94 01576 (SYSTEMIX INC) 20 January 1994 see the whole document ---	1,2,9-16
A	WO,A,93 15201 (NEW ENGLAND DEACONESS HOSPITAL) 5 August 1993 see the whole document ---	1-16
A	WO,A,92 14748 (AMERICAN CYANAMID CO) 3 September 1992 see the whole document -----	1-16

# INTERNATIONAL SEARCH REPORT

Intern. application No.

PCT/GB95/01213

## Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.: 10-13  
because they relate to subject matter not required to be searched by this Authority, namely:  
Remark: Although claims 10-13 are directed to a method of treatment of the human/animal body as well as diagnostic methods (Rule 39.1(iv) PCT) the search has been carried out and based on the alleged effects of the compound/composition.
2. ☐ Claims Nos.:  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☐ Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.



# INTERNATIONAL SEARCH REPORT

International Application No  
PCT/GB 95/01213

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
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WO-A-9315201	05-08-93	AU-B- 3482493 CA-A- 2128722 EP-A- 0624192 JP-T- 7504813	01-09-93 05-08-93 17-11-94 01-06-95
WO-A-9214748	03-09-92	EP-A- 0536350	14-04-93

